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Sequence Listing was accepted.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2008; month=3; day=19; hr=15; min=11; sec=5; ms=399;]

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Application No: 10542408 Version No: 2.0

Input Set:

Output Set:

Started: 2008-03-06 19:44:42.571
Finished: 2008-03-06 19:44:43.734
Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 163 ms
Total Warnings: 16
Total Errors: 2
No. of SeqIDs Defined: 22
Actual SeqID Count: 22

| Error code | Error Description |
|------------|---|
| W 213 | Artificial or Unknown found in <213> in SEQ ID (5) |
| W 213 | Artificial or Unknown found in <213> in SEQ ID (6) |
| W 213 | Artificial or Unknown found in <213> in SEQ ID (7) |
| W 213 | Artificial or Unknown found in <213> in SEQ ID (10) |
| W 213 | Artificial or Unknown found in <213> in SEQ ID (11) |
| W 213 | Artificial or Unknown found in <213> in SEQ ID (12) |
| W 213 | Artificial or Unknown found in <213> in SEQ ID (13) |
| W 213 | Artificial or Unknown found in <213> in SEQ ID (14) |
| W 213 | Artificial or Unknown found in <213> in SEQ ID (15) |
| W 213 | Artificial or Unknown found in <213> in SEQ ID (16) |
| W 213 | Artificial or Unknown found in <213> in SEQ ID (17) |
| W 213 | Artificial or Unknown found in <213> in SEQ ID (18) |
| W 213 | Artificial or Unknown found in <213> in SEQ ID (19) |
| W 213 | Artificial or Unknown found in <213> in SEQ ID (20) |
| W 213 | Artificial or Unknown found in <213> in SEQ ID (21) |
| E 257 | Invalid sequence data feature in <221> in SEQ ID (21) |
| W 213 | Artificial or Unknown found in <213> in SEQ ID (22) |
| E 257 | Invalid sequence data feature in <221> in SEQ ID (22) |

SEQUENCE LISTING

<110> Takeda Pharmaceutical Company Limited

<120> Novel Screening Method

<130> G05-0036

<140> 10542408

<141> 2008-03-06

<150> JP 2003-010001

<151> 2003-01-17

<150> JP 2003-104540

<151> 2003-04-08

<150> JP 2003-194497

<151> 2003-07-09

<150> JP 2003-329080

<151> 2003-09-19

<150> PCT/JP2004/000248

<151> 2004-01-15

<160> 22

<210> 1

<211> 361

<212> PRT

<213> Homo sapiens

<400> 1

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Leu Glu Gln Ala Asn Arg Thr Arg Phe Pro Phe Phe Ser Asp Val Lys
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Gly Asp His Arg Leu Val Leu Ala Ala Val Glu Thr Thr Val Leu Val
      35              40              45
Leu Ile Phe Ala Val Ser Leu Leu Gly Asn Val Cys Ala Leu Val Leu
      50              55              60
Val Ala Arg Arg Arg Arg Arg Gly Ala Thr Ala Cys Leu Val Leu Asn
      65              70              75              80
Leu Phe Cys Ala Asp Leu Leu Phe Ile Ser Ala Ile Pro Leu Val Leu
      85              90              95
Ala Val Arg Trp Thr Glu Ala Trp Leu Leu Gly Pro Val Ala Cys His
      100             105             110
Leu Leu Phe Tyr Val Met Thr Leu Ser Gly Ser Val Thr Ile Leu Thr
      115             120             125
Leu Ala Ala Val Ser Leu Glu Arg Met Val Cys Ile Val His Leu Gln
      130             135             140
Arg Gly Val Arg Gly Pro Gly Arg Arg Ala Arg Ala Val Leu Leu Ala
      145             150             155             160
Leu Ile Trp Gly Tyr Ser Ala Val Ala Ala Leu Pro Leu Cys Val Phe
      165             170             175
Phe Arg Val Val Pro Gln Arg Leu Pro Gly Ala Asp Gln Glu Ile Ser

```

| | | | | | |
|---|---|---------------------------------|-----|--|-----|
| | 180 | | 185 | | 190 |
| Ile Cys Thr | Leu Ile Trp Pro Thr | Ile Pro Gly Glu Ile Ser Trp Asp | | | |
| 195 | 200 | 205 | | | |
| Val Ser Phe | Val Thr Leu Asn Phe Leu Val Pro Gly Leu Val Ile Val | | | | |
| 210 | 215 | 220 | | | |
| Ile Ser Tyr Ser Lys Ile Leu Gln Ile Thr Lys Ala Ser Arg Lys Arg | | | | | |
| 225 | 230 | 235 | 240 | | |
| Leu Thr Val Ser Leu Ala Tyr Ser Glu Ser His Gln Ile Arg Val Ser | | | | | |
| 245 | 250 | 255 | | | |
| Gln Gln Asp Phe Arg Leu Phe Arg Thr Leu Phe Leu Leu Met Val Ser | | | | | |
| 260 | 265 | 270 | | | |
| Phe Phe Ile Met Trp Ser Pro Ile Ile Ile Thr Ile Leu Leu Ile Leu | | | | | |
| 275 | 280 | 285 | | | |
| Ile Gln Asn Phe Lys Gln Asp Leu Val Ile Trp Pro Ser Leu Phe Phe | | | | | |
| 290 | 295 | 300 | | | |
| Trp Val Val Ala Phe Thr Phe Ala Asn Ser Ala Leu Asn Pro Ile Leu | | | | | |
| 305 | 310 | 315 | 320 | | |
| Tyr Asn Met Thr Leu Cys Arg Asn Glu Trp Lys Lys Ile Phe Cys Cys | | | | | |
| 325 | 330 | 335 | | | |
| Phe Trp Phe Pro Glu Lys Gly Ala Ile Leu Thr Asp Thr Ser Val Lys | | | | | |
| 340 | 345 | 350 | | | |
| Arg Asn Asp Leu Ser Ile Ile Ser Gly | | | | | |
| 355 | 360 | | | | |

<210> 2
 <211> 1083
 <212> DNA
 <213> Homo sapiens

<400> 2

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| aaccgcaccc | gctttccctt | ctttctccgac | gtcaaggcg | accaccggct | ggtgctggcc | 120 |
| gcggtggaga | caaccgtgct | ggtgctcatc | tttgagtg | cgctgctggg | caacgtgtgc | 180 |
| gccttggtgc | tgggtggcgcg | ccgacgacgc | cgcggcgcga | ctgcctgcct | ggtactcaac | 240 |
| ctcttctgcg | cggacctgct | cttcacacgc | gctatccctc | tgggtgctggc | cgtgcgctgg | 300 |
| actgaggcct | ggctgctggg | ccccgttgcc | tgccacctgc | tcttctacgt | gatgaccctg | 360 |
| agcggcagcg | tcaccatcct | cacgctggcc | gcggtcagcc | tggagcgcct | ggtgtgcatc | 420 |
| gtgcacctgc | agcgcggcgt | gcggggctct | ggcgggcg | cgcgggcagt | gctgctggcg | 480 |
| ctcatctggg | gctattcggc | ggtcgccgct | ctgcctctct | gcgtcttctt | ccgagtcgtc | 540 |
| ccgcaacggc | tccccggcgc | cgaccaggaa | atttcgattt | gcacactgat | ttggcccacc | 600 |
| attctctggag | agatctcgtg | ggatgtctct | tttggtactt | tgaacttctt | ggtgccagga | 660 |
| ctgggtcattg | tgatcagtta | ctccaaaatt | ttacagatca | caaaggcatc | aaggaagagg | 720 |
| ctcacggtaa | gcctggccta | ctcgagagc | caccagatcc | gcgtgtccca | gcaggacttc | 780 |
| cggtcttctc | gcacctctt | cctcctcatg | gtctccttct | tcatcatgtg | gagccccatc | 840 |
| atcatcacca | tctcctcat | cctgatccag | aacttcaagc | aagacctggg | catctggccg | 900 |
| tccctcttct | tctgggtggg | ggccttcaca | tttgctaatt | cagccctaaa | ccccatcctc | 960 |
| tacaacatga | cactgtgcag | gaatgagtg | aagaaaattt | tttgctgctt | ctggttccca | 1020 |
| gaaaaggagg | ccattttaac | agacacatct | gtcaaaagaa | atgacttgtc | gattatttct | 1080 |
| ggc | | | | | | 1083 |

<210> 3
 <211> 361
 <212> PRT
 <213> Mus musculus

<400> 3

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| Met Ser Pro Glu Cys Ala Gln Thr Thr Gly Pro Gly Pro Ser His Thr |
|---|


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agcggcagcg tcacgaccc cactactggc gcggtcagcc tggagcgcac ggtgtgcatc 420
gtgcgcctcc ggcgcggttt gagcgccccc gggcgccgga ctcaggcggc actgctggct 480
ttcatatggg gttactcggc gctcgccgcg ctgcccctct gcatcttggt ccgcgtggtc 540
ccgcagcgcc ttcccgcgcg ggaccaggaa attccgattt gcacattgga ttggcccaac 600
cgcataggag aaatctcatg ggatgtgttt tttgtgactt tgaacttcct ggtgccggga 660
ctggtcattg tgatcagtta ctccaaaatt ttacagatca cgaaagcatc gcggaagagg 720
cttacgctga gcttggcata ctctgagagc caccagatcc gagtgtccca acaagactac 780
cgactcttcc gcacgctctt cctgctcatg gtttccttct tcatcatgtg gagtcccatc 840
atcatcacca tcctcctcat cttgatccaa aacttcgggc aggacctggg catctggcca 900
tcccttttct tctgggtggg ggccttcacg ttgccaact ctgcctaaa cccatactg 960
tacaacatgt cgctgttcag gaacgaatgg aggaagattt tttgctgctt cttttttcca 1020
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agc 1083

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<210> 5
<211> 20
<212> DNA
<213> Artificial Sequence

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<220>
<223> primer

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<400> 5
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```

<210> 6
<211> 20
<212> DNA
<213> Artificial Sequence

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<220>
<223> primer

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<400> 6
cgctgtggat gtctatttgc 20

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<210> 7
<211> 30
<212> DNA
<213> Artificial Sequence

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<220>
<223> primer

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<400> 7
agttcatttc cagtaccctc catcagtggc 30

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<210> 8
<211> 361
<212> PRT
<213> Rattus norvegicus

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<400> 8
Met Ser Pro Glu Cys Ala Gln Thr Thr Gly Pro Gly Pro Ser Arg Thr
          5              10              15
Pro Asp Gln Val Asn Arg Thr His Phe Pro Phe Phe Ser Asp Val Lys

```


| | | | | | | |
|-------------|-------------|------------|-------------|------------|------------|------|
| gtgcgcctgc | ggcgcggcctt | gagcggcccg | gggcggcgga | cgcaggcggc | gctgctggct | 480 |
| ttcatatggg | gttactcggc | gctcgcgcgc | ctgcccctct | gcattcttgt | ccgcgtggtc | 540 |
| ccgcagcgcc | ttcccggcgg | ggaccaggaa | attccgattt | gcacattgga | ttggcccaac | 600 |
| cgcataggag | aaatctcatg | ggatgtgttt | tttgtgactt | tgaacttcct | ggtaccagga | 660 |
| ctgggtcattg | tgatcagcta | ctccaagatt | ttacagatca | cgaaagcctc | gcggaagagg | 720 |
| cttacgctga | gcttggcata | ctccgagagc | caccagatcc | gagtgtccca | gcaggactac | 780 |
| cggctcttcc | gaacgctctt | cctgctcatg | gtttccttct | tcatcatgtg | gagtcccatc | 840 |
| atcatcacca | tcctcctcat | cttgatccag | aacttcgcgc | aggacctgg | tatctggccg | 900 |
| tcccttttct | tctgggtgg | ggccttcacg | tttgccaaact | ccgcctaaa | ccccattctg | 960 |
| tacaacatgt | cgtgttccag | gagcgagtgg | aggaagattt | tttgctgctt | ctttttccca | 1020 |
| gagaagggag | ccattttttac | agaaacgtct | atcaggcgaa | atgacttgtc | tgttatttcc | 1080 |
| acc | | | | | | 1083 |

<210> 10
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 <212> DNA
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<220>
 <223> primer

<400> 10
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<210> 11
 <211> 19
 <212> DNA
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<220>
 <223> primer

<400> 11
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<210> 12
 <211> 26
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> probe

<400> 12
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<210> 13
 <211> 33
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<220>
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<400> 13
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<210> 14
 <211> 33
 <212> DNA
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 <220>
 <223> primer

 <400> 14
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 <210> 15
 <211> 23
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 <220>
 <223> primer

 <400> 15
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 <210> 16
 <211> 24
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 <220>
 <223> primer

 <400> 16
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 <210> 17
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 <400> 17
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 <210> 18
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 <220>
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 <400> 18
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 <210> 19
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<212> DNA
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 <220>
 <223> primer

 <400> 19
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 <210> 20
 <211> 26
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 <220>
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 <400> 20
 caactccgcc ctaaacccca ttctgt 26

 <210> 21
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 <221> misc_DNA
 <222> (20)..(21)
 <223> n stands for deoxy thymidine

 <400> 21
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 <210> 22
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 <213> Artificial Sequence

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 <223> base sequence of the antisense strand of siRNA ml4i561

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 <221> misc_DNA
 <222> (1)..(2)
 <223> n stands for deoxy thymidine

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